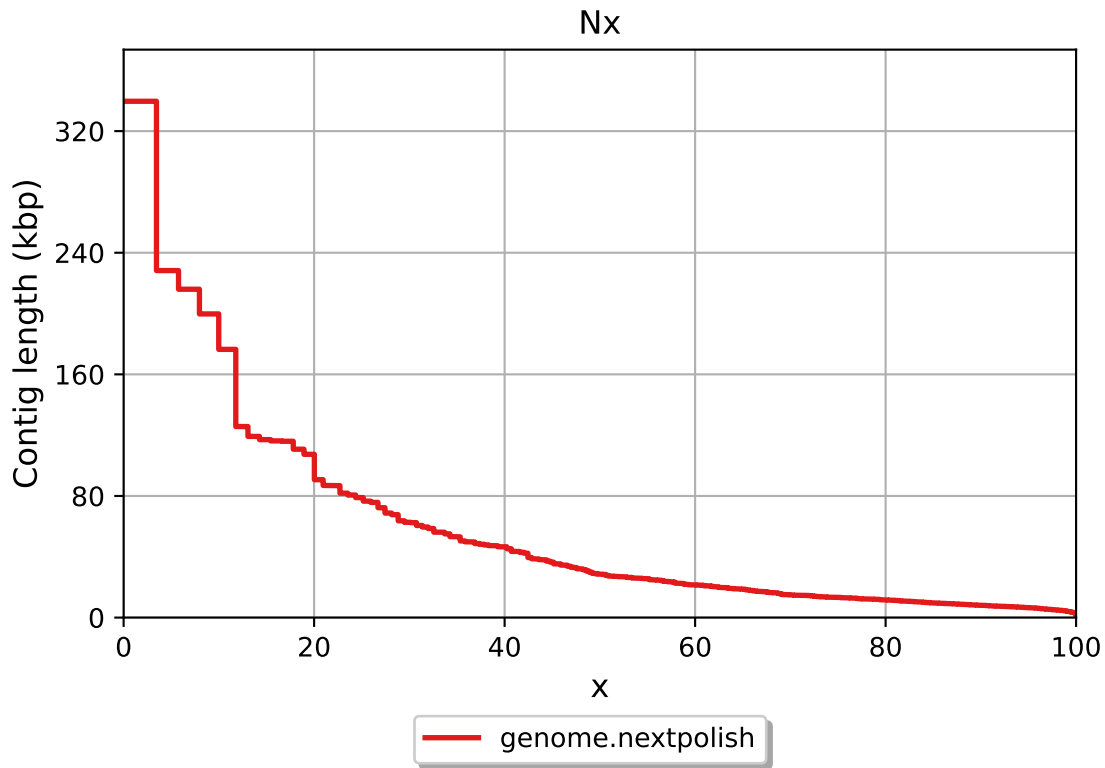


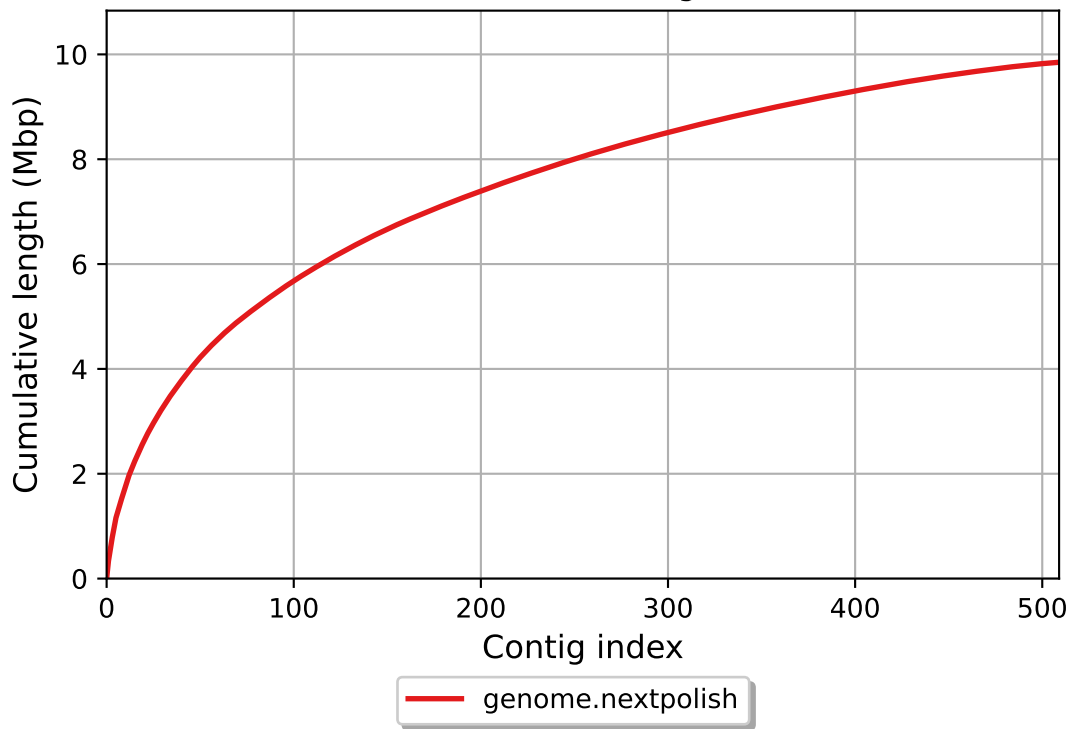
Report

	genome.nextpolish
# contigs (>= 0 bp)	509
# contigs (>= 1000 bp)	509
# contigs (>= 5000 bp)	460
# contigs (>= 10000 bp)	276
# contigs (>= 25000 bp)	90
# contigs (>= 50000 bp)	35
Total length (>= 0 bp)	9849616
Total length (>= 1000 bp)	9849616
Total length (>= 5000 bp)	9653845
Total length (>= 10000 bp)	8279945
Total length (>= 25000 bp)	5436811
Total length (>= 50000 bp)	3531143
# contigs	509
Largest contig	339633
Total length	9849616
GC (%)	48.44
N50	28652
N90	8071
auN	62547.7
L50	71
L90	341
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	27 + 9 part

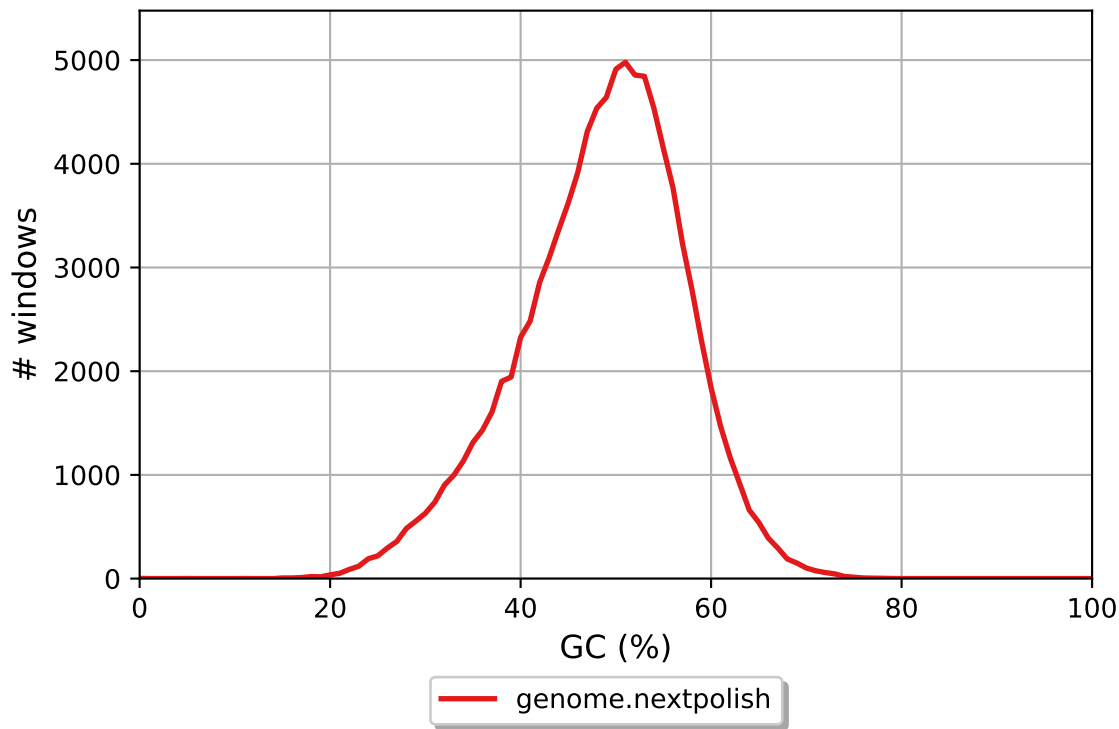
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



GC content



genome.nextpolish GC content

